52. The method of claim 30, wherein the amino acid sequence of said polypeptide is at least 80% identical to the corresponding region of a mature, native human apoE polypeptide, beginning at amino acid residue 1.

53. The method of claim 30, wherein said polypeptide has a signal peptide operably linked to said region of said mature apoE.

54. The method of claim 30, wherein said polypeptide consists of between 150 and 215 amino acids.

55. The method of claim 30, wherein said polypeptide consists of 203 amino acids.

56. The method of claim 30, wherein said nucleic acid encodes residues 1-203 of an apoE preprotein of any one of SEQ ID Nos. 14-19.

57. The method of claim 30, wherein said polypeptide consists of 220 amino acids.

58. The method of claim 30, wherein said nucleic acid encodes residues 1-220 of an apoE preprotein of any one of SEQ ID Nos. 14-19.

59. The method of claim 30, wherein said polypeptide consists of 247 amino acids.

60. The method of claim 30, wherein said nucleic acid encodes residues 1-247 of an apoE preprotein of any one of SEQ ID Nos. 14-19.

61. The method of claim 30, wherein said polypeptide consists of 277 amino acids.